

Figure 1

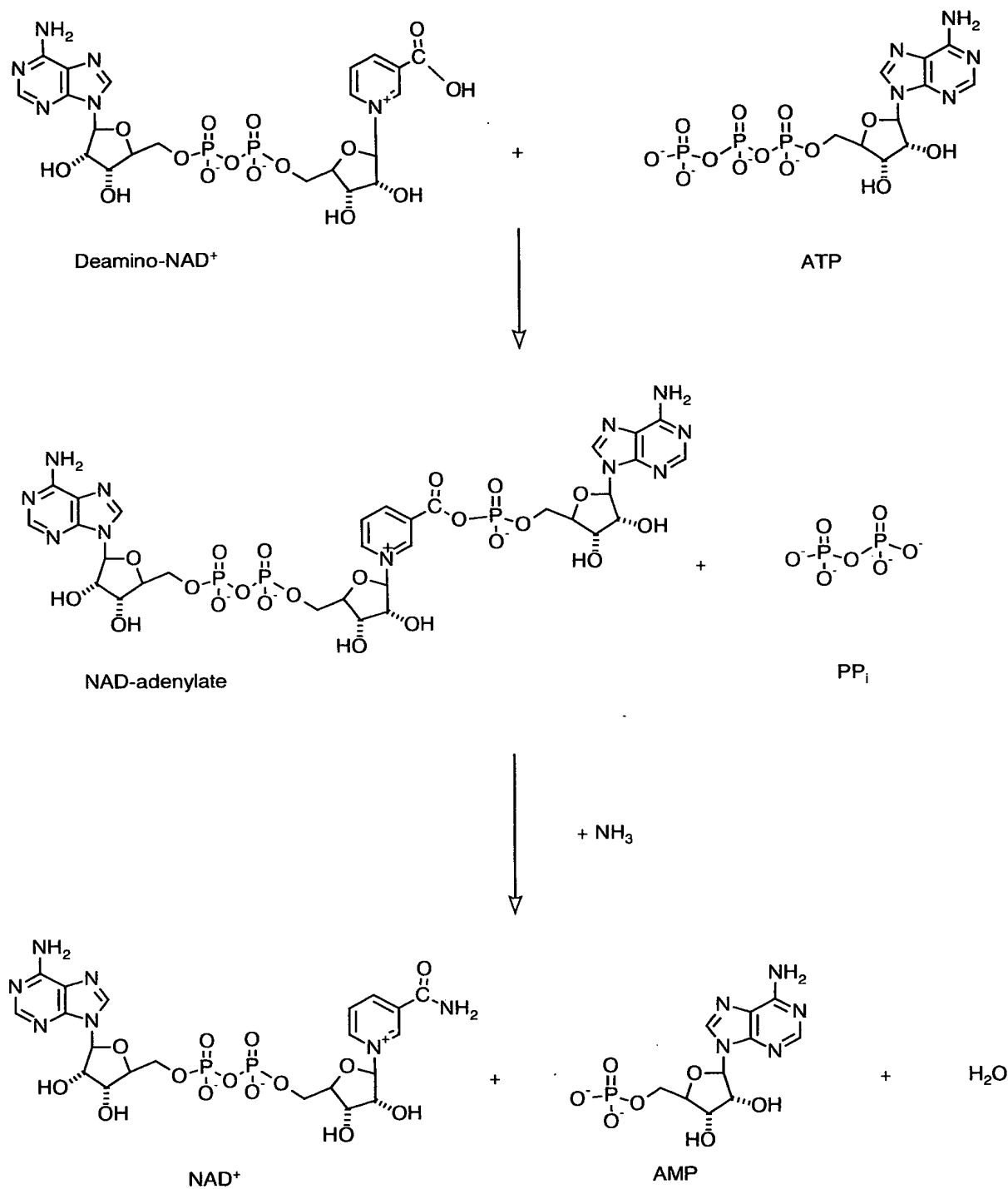
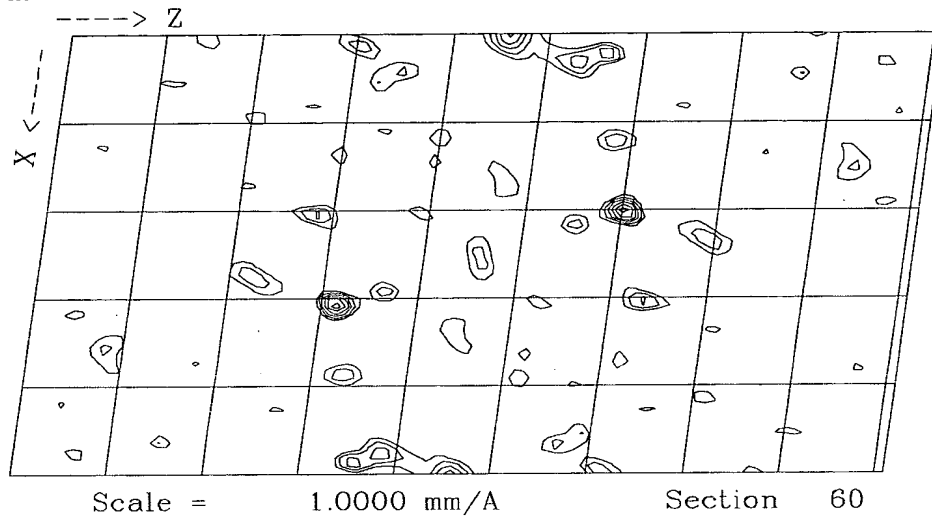


Figure 2

a.



b.

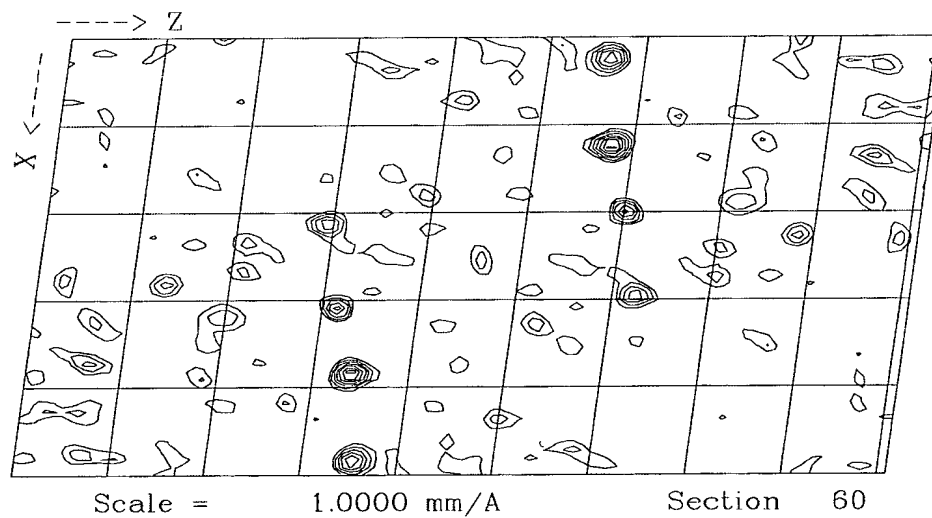
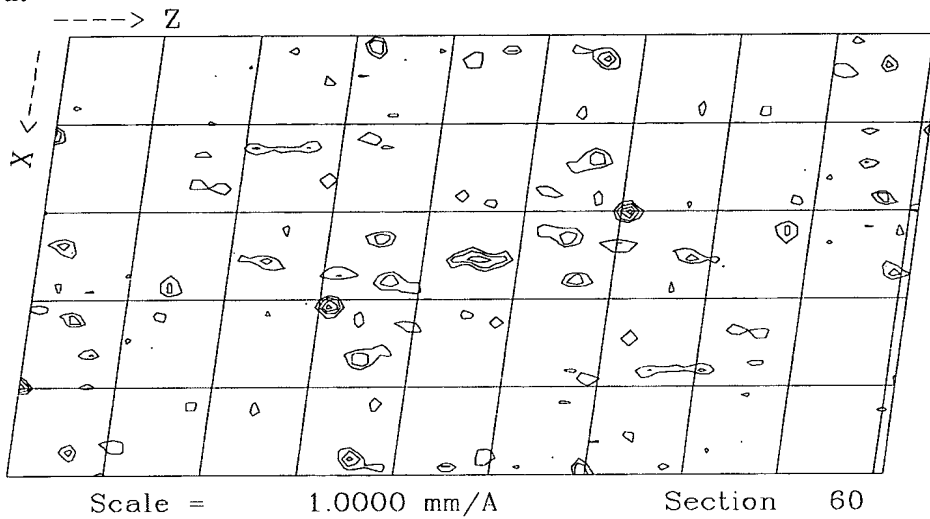


Figure 3

a.



b.

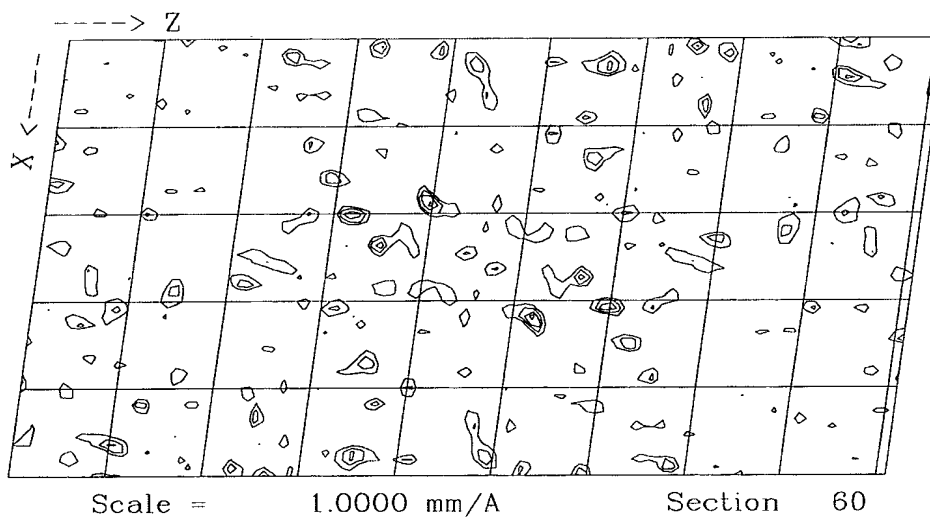
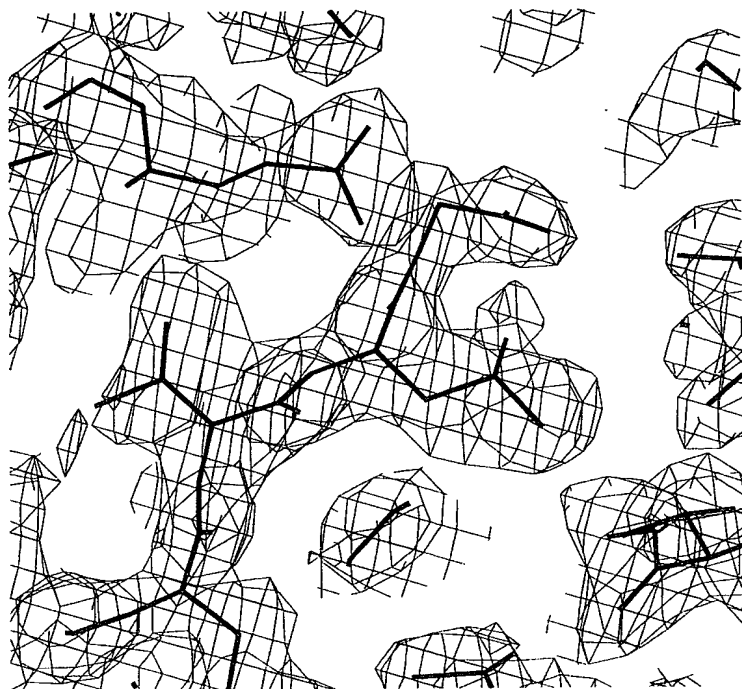


Figure 4

a.



b.

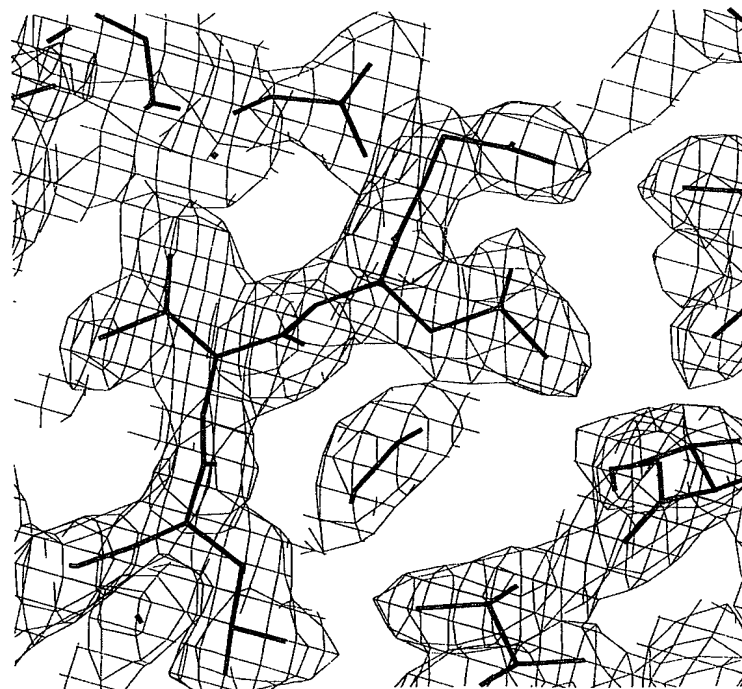


Figure 4a: 6315.N

Figure 5

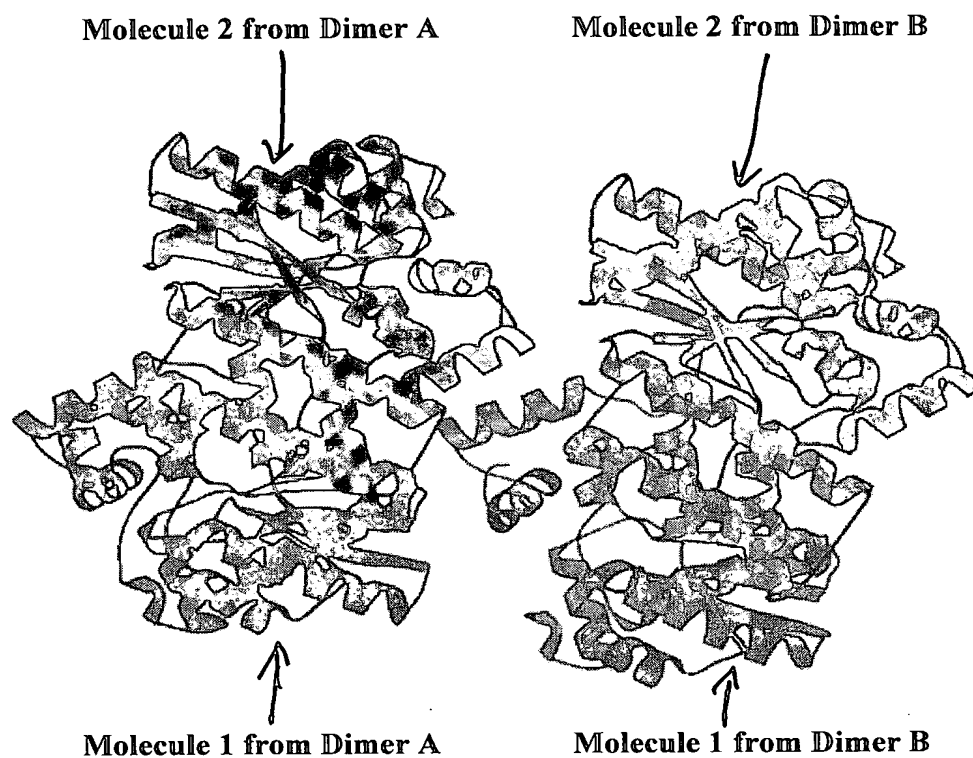


FIG. 5

### Figure 6

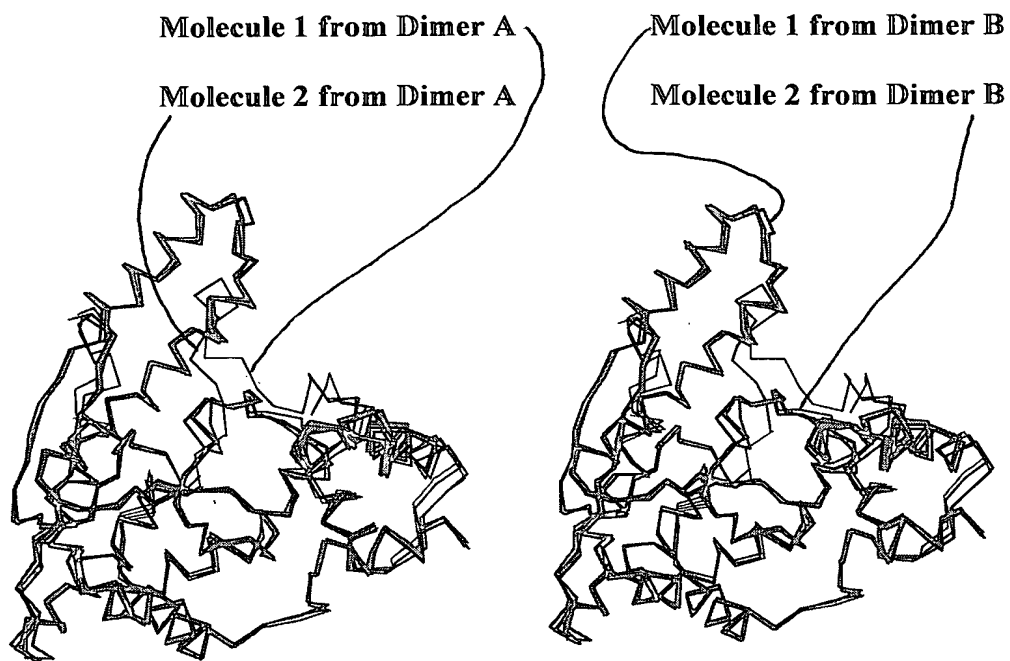
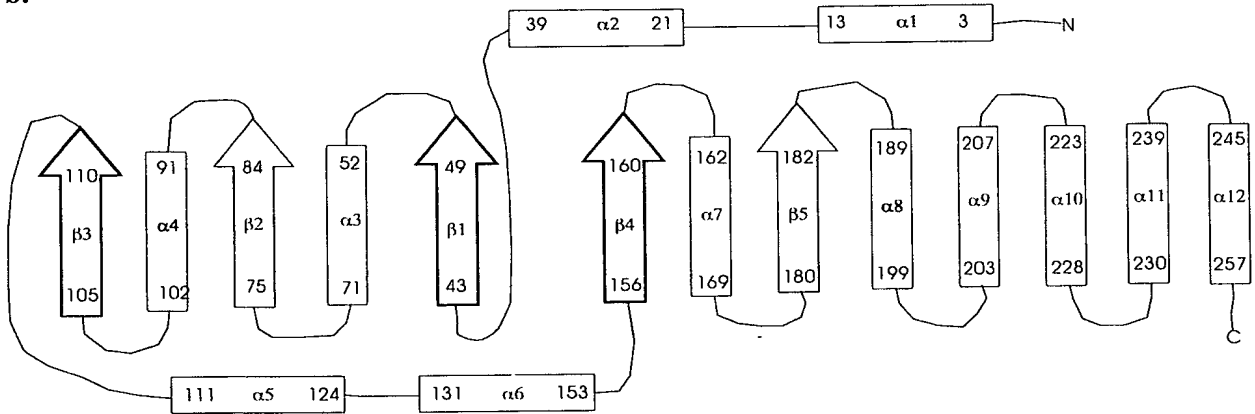


Figure 7

a.



b.



c.

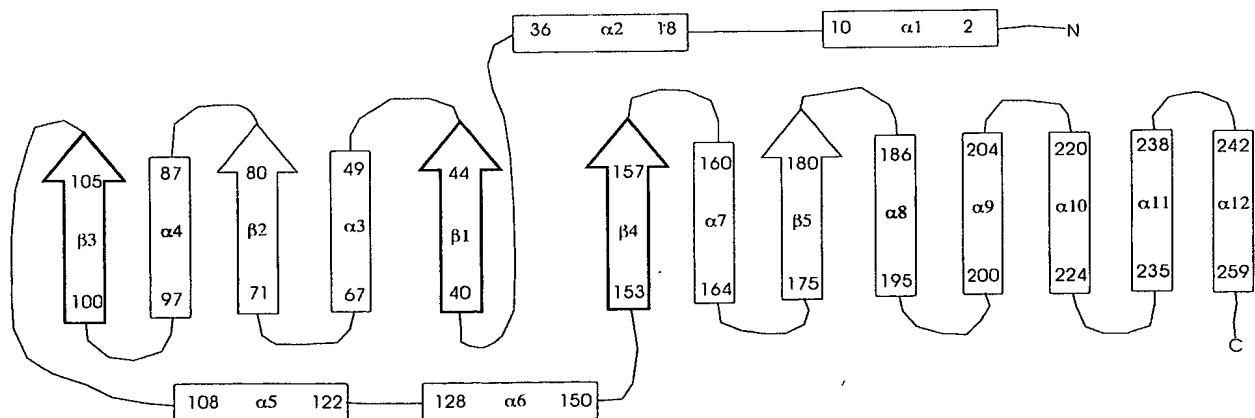


Figure 8



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### Figure 9

		1		50
nade_bs	(1)	---	SMQEKIMRELHVKPSIDPKQEIEDRVNFLKQYVK-KTGAKGFVLGIS	
nade_sa	(1)		MGSKEODVIVQEMKVKKRIDSAEEIMELKQFIKKNYVQSHSFIKSLVLGIS	
		51		100
nade_bs	(47)		GGQDSTLAGRLAQLAVESTREEGGDAQFIAVRLPHGTOQDEDDAQLALKEF	
nade_sa	(51)		GGQDSTLVGKLVMQSVNELREEGIDCTFIAVKLPVGVQKDADEVEQALRF	
		101		150
nade_bs	(97)		IKPKDSWKFDIKSTVSAFSDQYQOETGDQLTDFNKGNVKARTRMIAQYAI	
nade_sa	(101)		IEPDEIVTVNIKPAVDQS-VQSLKEAGIVLTDFOKGNEKARERMKVQESI	
		151		200
nade_bs	(147)		GGQEGLEVLTGDHAAEAVTGFFTKYGDGGADLPLTGLTKRQGRITLLKEL	
nade_sa	(150)		ASNRQGLVVGTDHSAENLTGFYTKYGDGAADTAPITFGLNKRQGRQLLAYL	
		201		250
nade_bs	(197)		GAPERLYLKEPTADLLDEKPOQSDTELGISYDEIDDYLEGKEVSAKVSE	
nade_sa	(200)		GAPKELYEKTPTADLEDDKPQLPDEDALGVITYEADINYLEGKPVITPEEQK	
		251		284
nade_bs	(247)		ALEKRYSMTEHKRQPPASMEDDWWK-----	
nade_sa	(250)		VLENHYIRNAHKRELAYLRYT-WPKSRSHHHHHH	

[illegible]

Figure 10

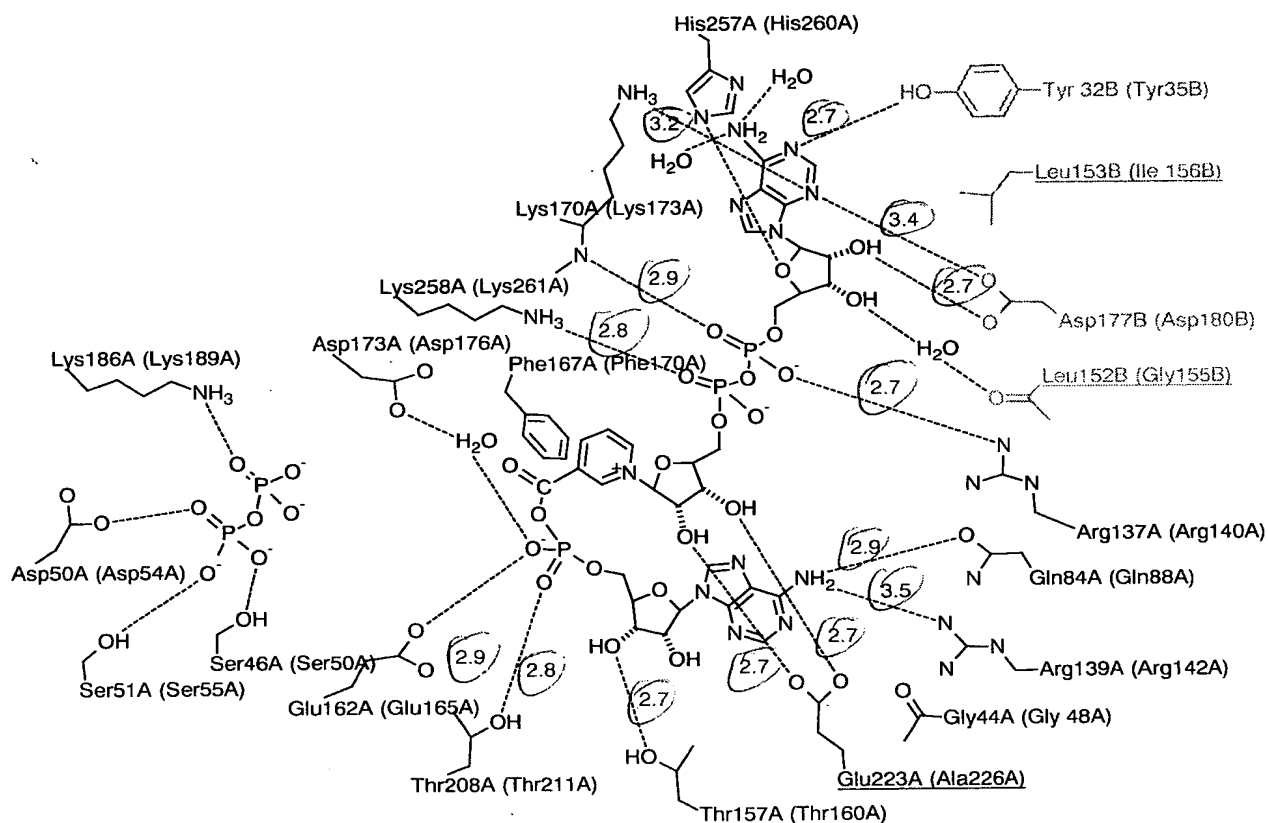


FIG. 10